OIPE

#5

RAW SEQUENCE LISTING
PATENT APPLICATION:  US/09/943,115A

DATE: 05/07/2002
TIME: 14:51:46

Input Set : D:\52459-20021.txt
Output Set: N:\CRF3\05072002\I943115A.raw

ENTERED

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 4 <110> APPLICANT: SEQUENOM, Inc.
 5        Risinger, Carl
 6        Andersson, Maria
 7        Lewander, Tommy
 8        Olaisson, Erik
10 <120> TITLE OF INVENTION: DETECTION OF CYP3A4 AND CYP2C9
11        POLYMORPHISMS
13 <130> FILE REFERENCE: 52459-20021.00
15 <140> CURRENT APPLICATION NUMBER: US 09/943,115A
16 <141> CURRENT FILING DATE: 2001-08-30
18 <150> PRIOR APPLICATION NUMBER: UK 0021286.0
19 <151> PRIOR FILING DATE: 2000-08-30
21 <160> NUMBER OF SEQ ID NOS: 73
23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 1345
27 <212> TYPE: DNA
28 <213> ORGANISM: Homo sapiens
30 <400> SEQUENCE: 1
31 ctgcagtgac cactgcccca tcattgctgg ctgaggtggt tggggtccat ctggctatct      60
32 gggcagctgt tctcttctct cctttctctc ctgtttccag acatgcagta tttccagaga     120
33 gaaggggcca ctctttggca aagaacctgt ctaacttgct atctatggca ggacctttga     180
34 agggttcaca ggaagcagca caaattgata ctattccacc aagccatcag ctccatctca     240
35 tccatgccct gtctctcctt taggggtccc cttgccaaca gaatcacaga ggaccagcct     300
36 gaaagtgcag agacagcagc tgaggcacag ccaagagctc tggctgtatt aatgacctaa     360
37 gaagtcacca gaaagtcaga aggatgcata gcagaggccc agcaatctca gctaagtcaa     420
38 ctccaccagc ctttctagtt gcccactgtg tgtacagcac sctggtaggg accagagcca     480
39 tgacagggaa taagactaga ctatgccctt gaggagctca cctctgttca gggaaacagg     540
40 cgtggaaaca caatggtggt aaagaggaaa gaggacaata ggattgcatg aaggggatgg     600
41 aaagtgccca ggggaggaaa tggttacatc tgtgtgagga gtttggtgag gaaagactct     660
42 aagagaaggc tctgtctgtc tgggtttgga aggatgtgta ggagtcttct agggggcaca     720
43 ggcacactcc aggcataggt aaagatctgt aggtgtggct tgttgggatg aatttcaagt     780
44 attttggaat gaggacagcc atagagacaa gggcargaga gaggcgattt aatagatttt     840
45 atgccaatgg ctccacttga gtttctgata agaacccaga acccttggac tccccagtaa     900
46 cattgattga gttgtttatg atacctcata gaatatgaac tcaaaggagg tcagtgagtg     960
47 gtgtgtgtgt gattctttgc caacttccaa ggtggagaag cctcttccaa ctgcaggcag    1020
48 agcacaggtg gccctgctac tggctgcagc tccagccctg cctccttctc tagcatataa    1080
49 acaatccaac agcctcactg aatcactgct gtgcagggca ggaaagctcc atgcacatag    1140
50 cccagcaaag agcaacacag agctgaaagg aagactcaga ggagagagat aagtaaggaa    1200
51 agtagtgatg gctctcatcc cagacttggc catggaaacc tggcttctcc tggctgtcag    1260
52 cctggtgctc ctctatctgt gagtaactgt tcaggctcct cttctctgtt tcttggactt    1320
53 ggggtcgtaa tcaggcctct ctttt                                          1345
55 <210> SEQ ID NO: 2
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56 <211> LENGTH: 19
57 <212> TYPE: DNA
58 <213> ORGANISM: Artificial Sequence
60 <220> FEATURE:
61 <223> OTHER INFORMATION: Oligonucleotide of CYP3A4 region
63 <400> SEQUENCE: 2
64 acaagggcaa gagagaggc                                            19
66 <210> SEQ ID NO: 3
67 <211> LENGTH: 19
68 <212> TYPE: DNA
69 <213> ORGANISM: Artificial Sequence
71 <220> FEATURE:
72 <223> OTHER INFORMATION: Oligonucleotide of CYP3A4 region
74 <400> SEQUENCE: 3
75 acaagggcag gagagaggc                                            19
77 <210> SEQ ID NO: 4
78 <211> LENGTH: 10
79 <212> TYPE: DNA
80 <213> ORGANISM: Artificial Sequence
82 <220> FEATURE:
83 <223> OTHER INFORMATION: Oligonucelotide of CYP3A4 region
85 <400> SEQUENCE: 4
86 agggcaagag                                                      10
88 <210> SEQ ID NO: 5
89 <211> LENGTH: 10
90 <212> TYPE: DNA
91 <213> ORGANISM: Artificial Sequence
93 <220> FEATURE:
94 <223> OTHER INFORMATION: Oligonucleotide of CYP3A4 region
96 <400> SEQUENCE: 5
97 agggcaggag                                                      10
99 <210> SEQ ID NO: 6
100 <211> LENGTH: 2438
101 <212> TYPE: DNA
102 <213> ORGANISM: Homo sapiens
104 <400> SEQUENCE: 6
105 gatctcagat atcccttcta tctacacatt atctataatt ctttctttct gtaaactgaa     60
106 aggtcctaga aggagccgca gctcagcagg agagaggagg agctgagctg ggacccctac    120
107 ctcctgagga atgaaatgat tattataaag acagcaaccg agcttatttt acccaaaata    180
108 aggtagtata tttctgttag agtttagagt ttcatgagtc agggaccaag ttattgcttt    240
109 tctttgccct gtataaaggc ttctccaagg cctttgactt acctaagtac taaatgttat    300
110 aaaaccaaac tcttctgacc tctcaatcta gtcaactggg gctgtaatta ttaatgaaat    360
111 taatgtttat tttgaaaata atttactaga ctgaattacg aaatcctgaa tcattgtaca    420
112 ctatcagtaa atattggtgg acccaactga actgaatgtt ttgcttgaaa tgaaaccttt    480
113 gagatgcagg gcttatgggt tctagtccca gctctagcac tagcagacag catgttcttg    540
114 gctaagatac tgaatcttca aggctcagct tcctcattcc ggaaatgggt caatttatt    600
115 gtaagcagag gtaattgaga gattcaaaag ggacatgagg tgtaacaatt ctctgtaaat    660
116 tgttagaatc cctgttaaaa atgaccagta aagctttgtg caactgtgtc ttgacataac    720
117 tttattttc ttaataaaag aaatggaaat aacctcacta gggaatttag aacaaatatg    780
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118 atgatatctt taaagaaaat ggctttgcac aagtattgac attaatgatc tagtaaagtg    840
119 tatctttcta gttgtattta gatcctcaac tcagtatgtc agctcctgtt aaggtctata    900
120 cattgtggtg gttctgtgct gtgggtccat ttagtgattt ccctacctcc catcttytat    960
121 tgcatccaca actgtggttc tgtccataat ttcctttgct ttctgtgcat tattacatca   1020
122 tatctgaaaa tgagaaacca aaaacaatrg aaagcagcca tgtctggagg tgactggggg   1080
123 gtcgagaagc cctagtttct caaaccctta gcaccaaatt tttccctcag ttacactgag   1140
124 cgtttcactt ctgcagtgat ggaraaggga gatcccttat ttcttctcat gagcatctct   1200
125 ggtgctgttt cccttagaga caaataaggg gttctattta atgtgaagcc tgttttatga   1260
126 acagaataaa tgtggtgtat attcagaata actaatgttt ggaagttgtt ttattttttgc   1320
127 taaaaattgt tctcaaggca gctctggtgt aagagataat acaccacgat gggcatcaga   1380
128 agacctcagc tcaaatccca gttctgccag ctatgagctg tgtggcacca acaggtgtcc   1440
129 tgttctccca gggtctccct tttcccattt gaaaaataaa aaataacaat tcctgccttc   1500
130 aggaattttt tttaggggggt ttaatkgtaa aggtgtttat atctgctaag gtaatttact   1560
131 tgatatatgt ttggttattt aagatatatg agttatgtta gctatttcat gtttaggctg   1620
132 ctgtattttt agtaggctat attaaatatt tgaaaggatt wmattataaa gaacaaagtc   1680
133 tcctaatctt tgatatagca ttgacatact ttttaaatat acaaggcata gaatatggcc   1740
134 atttctgtta aatcatatat tcccaactgg ttattaatct aagaattcag aattttgagt   1800
135 aattgctttt gcatcagatt atttacttca gtgctctcaa ttatgatggt gcattagaac   1860
136 catctgggtt aacatttgtt ttttattacc aatacctagg ctccaaccaa gtacagtgaa   1920
137 actggaatgt acagagtgga caatggaacg aaggagaaca agaccaaagg acattttatt   1980
138 tttatctgta tcagtgggtc aaagtccttt cagaaggagc atatagtgga cctaggtgat   2040
139 tggtcaattt atccatcaaa gaggcacaca ccgaattagc atggagtgtt ataaaaggct   2100
140 tggagtgcaa gctcatggtt gtcttaacaa gaagagaagg cttcaatgga ttctcttgtg   2160
141 gtccttgtgc tctgtctctc atgtttgctt ctcctttcac tctggagaca gagctctggg   2220
142 agaggaaaac tccctcctgg ccccactcct ctcccagtga ttggaaatat cctacagata   2280
143 ggtattaagg acatcagcaa atccttaacc aatgtaagta tgctccttca gtggcttgca   2340
144 aaaggtaagt aaattcacct gtattttta aataaagtgt atccctagag gtacatgtta   2400
145 caagaggtaa tggtaaagta aaatactttg aaaggctt                            2438
147 <210> SEQ ID NO: 7
148 <211> LENGTH: 20
149 <212> TYPE: DNA
150 <213> ORGANISM: Artificial Sequence
152 <220> FEATURE:
153 <223> OTHER INFORMATION: Primer
155 <400> SEQUENCE: 7
156 ccagcctgaa agtgcagaga                                                 20
158 <210> SEQ ID NO: 8
159 <211> LENGTH: 25
160 <212> TYPE: DNA
161 <213> ORGANISM: Artificial Sequence
163 <220> FEATURE:
164 <223> OTHER INFORMATION: Primer
166 <400> SEQUENCE: 8
167 tcttagagtc tttcctcacc aaact                                          25
169 <210> SEQ ID NO: 9
170 <211> LENGTH: 20
171 <212> TYPE: DNA
172 <213> ORGANISM: Artificial Sequence
174 <220> FEATURE:
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175 <223> OTHER INFORMATION: Primer
177 <400> SEQUENCE: 9
178 catgccctgt ctctccttta                                              20
180 <210> SEQ ID NO: 10
181 <211> LENGTH: 19
182 <212> TYPE: DNA
183 <213> ORGANISM: Artificial Sequence
185 <220> FEATURE:
186 <223> OTHER INFORMATION: Primer
188 <400> SEQUENCE: 10
189 ccatcccctt catgcaatc                                               19
191 <210> SEQ ID NO: 11
192 <211> LENGTH: 11
193 <212> TYPE: DNA
194 <213> ORGANISM: Artificial Sequence
196 <220> FEATURE:
197 <223> OTHER INFORMATION: Oligonucleotide representing the coding strand
198       complementary to the polymorphic site 461
200 <400> SEQUENCE: 11
201 agcaccctgg t                                                       11
203 <210> SEQ ID NO: 12
204 <211> LENGTH: 11
205 <212> TYPE: DNA
206 <213> ORGANISM: Artificial Sequence
208 <220> FEATURE:
209 <223> OTHER INFORMATION: Oligonucleotide representing the coding strand
210       complementary to the polymorphic site 461
212 <400> SEQUENCE: 12
213 agcacgctgg t                                                       11
215 <210> SEQ ID NO: 13
216 <211> LENGTH: 11
217 <212> TYPE: DNA
218 <213> ORGANISM: Artificial Sequence
220 <220> FEATURE:
221 <223> OTHER INFORMATION: Oligonucleotide representing the non-coding strand
222       complementary to the polymorphic site 461
224 <400> SEQUENCE: 13
225 accagggtgc t                                                       11
227 <210> SEQ ID NO: 14
228 <211> LENGTH: 11
229 <212> TYPE: DNA
230 <213> ORGANISM: Artificial Sequence
232 <220> FEATURE:
233 <223> OTHER INFORMATION: Oligonucleotide representing the non-coding strand
234       complementary to the polymorphic site 461
236 <400> SEQUENCE: 14
237 accagcgtgc t                                                       11
239 <210> SEQ ID NO: 15
240 <211> LENGTH: 11
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241 <212> TYPE: DNA
242 <213> ORGANISM: Artificial Sequence
244 <220> FEATURE:
245 <223> OTHER INFORMATION: Oligonucleotide of the novel polymorphic site 461
246       on the coding strand
248 <400> SEQUENCE: 15
249 gtgtgtacag c                                                              11
251 <210> SEQ ID NO: 16
252 <211> LENGTH: 11
253 <212> TYPE: DNA
254 <213> ORGANISM: Artificial Sequence
256 <220> FEATURE:
257 <223> OTHER INFORMATION: Oligonucleotide of the novel polymorphic site 461
258       on the non-coding strand
260 <400> SEQUENCE: 16
261 gctgtacaca c                                                              11
263 <210> SEQ ID NO: 17
264 <211> LENGTH: 11
265 <212> TYPE: DNA
266 <213> ORGANISM: Artificial Sequence
268 <220> FEATURE:
269 <223> OTHER INFORMATION: Oligonucleotide of the novel polymorphic site 461
270       on the non-coding strand
272 <400> SEQUENCE: 17
273 tggtccctac c                                                              11
275 <210> SEQ ID NO: 18
276 <211> LENGTH: 11
277 <212> TYPE: DNA
278 <213> ORGANISM: Artificial Sequence
280 <220> FEATURE:
281 <223> OTHER INFORMATION: Oligonucleotide of the novel polymorphic site 461
282       on the coding strand
284 <400> SEQUENCE: 18
285 ggtagggacc a                                                              11
287 <210> SEQ ID NO: 19
288 <211> LENGTH: 25
289 <212> TYPE: DNA
290 <213> ORGANISM: Artificial Sequence
292 <220> FEATURE:
293 <223> OTHER INFORMATION: Primer
295 <400> SEQUENCE: 19
296 cactagggaa tttagaacaa atatg                                               25
298 <210> SEQ ID NO: 20
299 <211> LENGTH: 23
300 <212> TYPE: DNA
301 <213> ORGANISM: Artificial Sequence
303 <220> FEATURE:
304 <223> OTHER INFORMATION: Primer
306 <400> SEQUENCE: 20
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**VERIFICATION SUMMARY**                 DATE: 05/07/2002
PATENT APPLICATION:   US/09/943,115A     TIME: 14:51:47

Input Set  : D:\52459-20021.txt
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